## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

| Application Serial Number: | 10/574,398 |
|----------------------------|------------|
| Source:                    | TFWP       |
| Date Processed by STIC:    | 04/26/2006 |

## ENTERED



**IFWP** 

RAW SEQUENCE LISTING DATE: 04/26/2006
PATENT APPLICATION: US/10/574,398 TIME: 07:32:52

Input Set : F:\5176.ST25.txt

Output Set: N:\CRF4\04262006\J574398.raw

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3 <110> APPLICANT: Bayer Pharamceuticals Corporation
             Pauloski, Nicole
             Taylor, Ian
     5
             Bigwood, Douglas
     6
     8 <120> TITLE OF INVENTION: GENE EXPRESSION PROFILES AND METHODS OF USE
    10 <130> FILE REFERENCE: 5176
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/574,398
                                                                (Pg-6)
C--> 12 <141> CURRENT FILING DATE: 2006-03-31
    12 <150> PRIOR APPLICATION NUMBER: PCT/US04/34163
     13 <151> PRIOR FILING DATE: 2004-10-01
    15 <150> PRIOR APPLICATION NUMBER: US 60/508,355
    16 <151> PRIOR FILING DATE: 2003-10-03
    18 <160> NUMBER OF SEQ ID NOS: 400
    20 <170> SOFTWARE: PatentIn version 3.3
     22 <210> SEQ ID NO: 1
     23 <211> LENGTH: 1863
     24 <212> TYPE: DNA
     25 <213> ORGANISM: Homo sapiens
     27 <400> SEQUENCE: 1
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                                                                              120
     30 tettteteca agtggtteet ettgagggga geatttetge tggetecagg aetttggeea
                                                                              180
     32 totataaago ttggcaatga gaaataagaa aattotoaag gaggacgago tottgagtga
                                                                              240
     34 gacccaacaa gctgcttttc accaaattgc aatggagcct ttcgaaatca atgttccaaa
     36 gcccaagagg agaaatgggg tgaacttctc cctagctgtg gtggtcatct acctgatcct
                                                                              300
     38 gctcaccgct ggcgctgggc tgctggtggt ccaagttctg aatctgcagg cgcggctccg
                                                                              360
                                                                              420
     40 ggtcctggag atgtatttcc tcaatgacac tctggcggct gaggacagcc cgtccttctc
     42 cttgctgcag tcagcacacc ctggagaaca cctggctcag ggtgcatcga ggctgcaagt
                                                                              480
                                                                              540
     44 cctgcaggcc caactcacct gggtccgcgt cagccatgag cacttgctgc agcgggtaga
                                                                              600
     46 caacttcact cagaacccag ggatgttcag aatcaaaggt gaacaaggcg ccccaggtct
                                                                              660
     48 tcaaggccac aagggggcca tgggcatgcc tggtgcccct ggcccgccgg gaccacctgc
                                                                              720
     50 tgagaaggga gccaaggggg ctatgggacg agatggagca acaggcccct cgggacccca
     52 aggcccaccg ggagtcaagg gagaggcggg cctccaagga ccccagggtg ctccagggaa
                                                                              780
                                                                              840
    54 gcaaggagcc actggcaccc caggacccca aggagagaag ggcagcaaag gcgatggggg
                                                                              900
     56 tctcattggc ccaaaagggg aaactggaac taagggagag aaaggagacc tgggtctccc
     58 aggaagcaaa ggggacaggg gcatgaaagg agatgcaggg gtcatggggc ctcctggagc
                                                                              960
                                                                             1020
     60 ccaggggagt aaaggtgact tcgggaggcc aggcccacca ggtttggctg gttttcctgg
                                                                             1080
     62 agetaaagga gatcaaggac aacetggact geagggtgtt cegggeeete etggtgeagt
     64 gggacaccca ggtgccaagg gtgagcctgg cagtgctggc tcccctgggc gagcaggact
                                                                             1140
     66 tccagggagc cccgggagtc caggagccac aggcctgaaa ggaagcaaag gggacacagg
                                                                             1200
                                                                             1260
     68 acttcaagga cagcaaggaa gaaaaggaga atcaggagtt ccaggccctg caggtgtgaa
     70 gggagaacag gggagcccag ggctggcagg tcccaaggga gcccctggac aagctggcca
                                                                             1320
     72 gaagggagac cagggagtga aaggatette tggggagcaa ggagtaaagg gagaaaaagg
                                                                             1380
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74 tgaaagaggt gaaaactcag tgtccgtcag gattgtcggc agtagtaacc gaggccgggc

1440

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Input Set : F:\5176.ST25.txt

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76 tqaaqtttac tacagtggta cctgggggac aatttgcgat gacgagtggc aaaattctga
78 tgccattgtc ttctgccgca tgctgggtta ctccaaagga agggccctgt acaaagtggg
                                                                     1560
80 agctggcact gggcagatct ggctggataa tgttcagtgt cggggcacgg agagtaccct
                                                                     1620
82 gtggagctgc accaagaata gctggggcca tcatgactgc agccacgagg aggacgcagg
                                                                     1680
84 cgtggagtgc agcgtctgac ccggaaaccc tttcacttct ctgctcccga ggtgtcctcg
                                                                     1740
86 qqctcatatq tqqqaaqqca gaggatctct gaggagttcc ctggggacaa ctgagcagcc
                                                                     1800
1860
                                                                     1863
90 aaa
93 <210> SEQ ID NO: 2
94 <211> LENGTH: 520
95 <212> TYPE: PRT
96 <213> ORGANISM: homo sapiens
98 <400> SEQUENCE: 2
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101 1
104 Gln Gln Ala Ala Phe His Gln Ile Ala Met Glu Pro Phe Glu Ile Asn
                                   25
108 Val Pro Lys Pro Lys Arg Arg Asn Gly Val Asn Phe Ser Leu Ala Val
                               40
112 Val Val Ile Tyr Leu Ile Leu Leu Thr Ala Gly Ala Gly Leu Leu Val
                           55
116 Val Gln Val Leu Asn Leu Gln Ala Arg Leu Arg Val Leu Glu Met Tyr
                       70
120 Phe Leu Asn Asp Thr Leu Ala Ala Glu Asp Ser Pro Ser Phe Ser Leu
                                       90
                   85
124 Leu Gln Ser Ala His Pro Gly Glu His Leu Ala Gln Gly Ala Ser Arg
               100
                                   105
128 Leu Gln Val Leu Gln Ala Gln Leu Thr Trp Val Arg Val Ser His Glu
                               120
132 His Leu Leu Gln Arg Val Asp Asn Phe Thr Gln Asn Pro Gly Met Phe
                           135
133
136 Arg Ile Lys Gly Glu Gln Gly Ala Pro Gly Leu Gln Gly His Lys Gly
                                           155
                       150
140 Ala Met Gly Met Pro Gly Ala Pro Gly Pro Pro Gly Pro Pro Ala Glu
                   165
                                       170
144 Lys Gly Ala Lys Gly Ala Met Gly Arg Asp Gly Ala Thr Gly Pro Ser
                                   185
148 Gly Pro Gln Gly Pro Pro Gly Val Lys Gly Glu Ala Gly Leu Gln Gly
                               200
           195
152 Pro Gln Gly Ala Pro Gly Lys Gln Gly Ala Thr Gly Thr Pro Gly Pro
                           215
                                               220
156 Gln Gly Glu Lys Gly Ser Lys Gly Asp Gly Gly Leu Ile Gly Pro Lys
                                           235
                       230
160 Gly Glu Thr Gly Thr Lys Gly Glu Lys Gly Asp Leu Gly Leu Pro Gly
164 Ser Lys Gly Asp Arg Gly Met Lys Gly Asp Ala Gly Val Met Gly Pro
                                   265
168 Pro Gly Ala Gln Gly Ser Lys Gly Asp Phe Gly Arg Pro Gly Pro Pro
                               280
169
           275
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PATENT APPLICATION: US/10/574,398 TIME: 07:32:52

Input Set : F:\5176.ST25.txt

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172 Gly Leu Ala Gly Phe Pro Gly Ala Lys Gly Asp Gln Gly Gln Pro Gly 295 176 Leu Gln Gly Val Pro Gly Pro Pro Gly Ala Val Gly His Pro Gly Ala 310 315 180 Lys Gly Glu Pro Gly Ser Ala Gly Ser Pro Gly Arg Ala Gly Leu Pro 325 184 Gly Ser Pro Gly Ser Pro Gly Ala Thr Gly Leu Lys Gly Ser Lys Gly 340 345 188 Asp Thr Gly Leu Gln Gly Gln Gly Arg Lys Gly Glu Ser Gly Val 189 355 360 192 Pro Gly Pro Ala Gly Val Lys Gly Glu Gln Gly Ser Pro Gly Leu Ala 375 196 Gly Pro Lys Gly Ala Pro Gly Gln Ala Gly Gln Lys Gly Asp Gln Gly 390 200 Val Lys Gly Ser Ser Gly Glu Gln Gly Val Lys Gly Glu Lys Gly Glu 405 410 204 Arg Gly Glu Asn Ser Val Ser Val Arg Ile Val Gly Ser Ser Asn Arg 420 425 208 Gly Arg Ala Glu Val Tyr Tyr Ser Gly Thr Trp Gly Thr Ile Cys Asp 440 212 Asp Glu Trp Gln Asn Ser Asp Ala Ile Val Phe Cys Arg Met Leu Gly 455 216 Tyr Ser Lys Gly Arg Ala Leu Tyr Lys Val Gly Ala Gly Thr Gly Gln 475 217 465 470 220 Ile Trp Leu Asp Asn Val Gln Cys Arg Gly Thr Glu Ser Thr Leu Trp 485 490 224 Ser Cys Thr Lys Asn Ser Trp Gly His His Asp Cys Ser His Glu Glu 228 Asp Ala Gly Val Glu Cys Ser Val 229 515 232 <210> SEQ ID NO: 3 233 <211> LENGTH: 816 234 <212> TYPE: DNA 235 <213> ORGANISM: Homo sapiens 238 <220> FEATURE: 239 <221> NAME/KEY: misc feature 240 <222> LOCATION: (293)..(293) 241 <223> OTHER INFORMATION: n is a, c, g, or t 243 <220> FEATURE: 244 <221> NAME/KEY: misc\_feature 245 <222> LOCATION: (509)..(509) 246 <223> OTHER INFORMATION: n is a, c, g, or t 248 <220> FEATURE: 249 <221> NAME/KEY: misc feature 250 <222> LOCATION: (534)..(534) 251 <223> OTHER INFORMATION: n is a, c, g, or t 253 <220> FEATURE: 254 <221> NAME/KEY: misc feature 255 <222> LOCATION: (624)..(624)

RAW SEQUENCE LISTING DATE: 04/26/2006 PATENT APPLICATION: US/10/574,398 TIME: 07:32:52

Input Set : F:\5176.ST25.txt

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    259 <221> NAME/KEY: misc_feature
    260 <222> LOCATION: (657)..(657)
    261 <223> OTHER INFORMATION: n is a, c, g, or t
    263 <220> FEATURE:
    264 <221> NAME/KEY: misc feature
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    268 <400> SEQUENCE: 3
    269 ttttqctaaa cattttttta aqtatqaqtc cttgtttaaa aagaaaagat taaaacagaa
                                                                                60
    271 aatattttct ataaataata catqtatttt ggttttagtg ctcccgccct aaggtttgaa
                                                                               120
    273 qtttactttt atccagtacc tttttcctcc atgatcacct ttttttctct ttcccctctc
                                                                               180
     275 ccactcgtgc acacgtgggg gtttctgcga gaattggcct tgctgcactg tgattggcga
                                                                               240
W--> 277 agacgtgaaa ctttttaaaa aaaatactta aaattggttc ttttgtttca ttntgtgtat
                                                                               300
     279 ttgaagtttt agttateete agaeteetet tetgetteee geageeaegt gaagaatgee
                                                                               360
     281 gtgacagatt tcagagccac gcccttccca ttctgctctg cagggtcctt gctgctctcc
                                                                               420
     283 catttgtaga aggcateete ggagateace teetegteat atagacaate aaaaaacate
                                                                               480
W--> 285 cgcagcacat tggcaggttg atcaagttnt actatcgatg cttgtagtgc atanagtgct
                                                                               540
                                                                               600
     287 tgcagttcct tctctgtatc tgagtctagg tacttgagta agatcggcac tctctgcttg
                                                                               660
W--> 289 ataacagcag tgtccactct gaangtagaa gagtcggcta taatagctgc tttacanaca
                                                                               720
W--> 291 gcagtcatta nagctctaac gaatgtagtg aactcatctg gatttcgtct agattagcct
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     293 ctacccagte aaagatetgt cateattege tttgetetea attatgagtt tetegagttg
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     298 <210> SEQ ID NO: 4
    299 <211> LENGTH: 1585
    300 <212> TYPE: PRT
    301 <213> ORGANISM: Homo sapiens
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    306 1
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    309 Ile Gln Pro Pro Arg Ala Thr Ile Pro Asn Ser Ser Pro Ser Ile Arg
                     20
                                         25
    313 Pro Gly Ala Gln Thr Pro Thr Ala Val Tyr Gln Ala Asn Gln His Ile
                                     40
    317 Met Met Val Asn His Leu Pro Met Pro Tyr Pro Val Pro Gln Gly Pro
     321 Gln Tyr Cys Ile Pro Gln Tyr Arg His Ser Gly Pro Pro Tyr Val Gly
                                                 75
                             70
    325 Pro Pro Gln Gln Tyr Pro Val Gln Pro Pro Gly Pro Gly Pro Phe Tyr
                                             90
    329 Pro Gly Pro Gly Pro Gly Asp Phe Pro Asn Ala Tyr Gly Thr Pro Phe
                                         105
    330
                     100
    333 Tyr Pro Ser Gln Pro Val Tyr Gln Ser Ala Pro Ile Ile Val Pro Thr
    334
                 115
                                     120
    337 Gln Gln Gln Pro Pro Pro Ala Lys Arg Glu Lys Lys Thr Ile Arg Ile
                                                      140
                                 135
     341 Arg Asp Pro Asn Gln Gly Gly Lys Asp Ile Thr Glu Glu Ile Met Ser
    342 145
                             150
                                                 155
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/574,398

DATE: 04/26/2006
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| 345<br>346 | Gly   | Gly        | Gly | Ser         | Arg<br>165 | Asn    | Pro        | Thr   | Pro        | Pro<br>170 | Ile  | Gly        | Arg  | Pro        | Thr<br>175 | Ser       |
|------------|-------|------------|-----|-------------|------------|--------|------------|-------|------------|------------|------|------------|------|------------|------------|-----------|
|            | Thr   | Pro        | Thr | Pro<br>180  |            | Gln    | Gln        | Leu   | Pro<br>185 |            | Gln  | Val        | Pro  | Glu<br>190 | His        | Ser       |
|            | Pro   | Val        | Val |             | Gly        | Thr    | Val        | Glu   |            | Ala        | His  | Leu        | Ala  |            | Ser        | Thr       |
| 354        |       |            | 195 |             |            |        | _          | 200   | _          |            |      |            | 205  | _          | _          | _         |
| 357<br>358 | Pro   | Val<br>210 | Thr | Ala         | Ala        | Ser    | Asp<br>215 | GIn   | Lys        | Gin        | GIu  | G1u<br>220 | Lys  | Pro        | Lys        | Pro       |
|            | Asp   |            | Val | Leu         | Lys        | Ser    |            | Ser   | Pro        | Val        | Leu  |            | Leu  | Val        | Leu        | Ser       |
| 362        | 225   |            |     |             |            | 230    |            |       |            |            | 235  |            |      |            |            | 240       |
|            | Gly   | Glu        | Lys | Lys         |            | Gln    | Glu        | Gly   | Gln        |            | Ser  | Glu        | Thr  | Thr        |            | Ile       |
| 366        | U = I | Sar        | Tla | Δla         | 245<br>Glu | T.e.ii | Pro        | I.e.i | Pro        | 250<br>Pro | Ser  | Pro        | Thr  | Thr        | 255<br>Val | Ser       |
| 370        | vai   | Ser        | 116 | 260         | GIU        | БСи    | 110        | Leu   | 265        | 110        | DCI  | 110        | 1111 | 270        | vai        | 502       |
| 373        | Ser   | Val        | Ala | Arg         | Ser        | Thr    | Ile        | Ala   | Ala        | Pro        | Thr  | Ser        | Ser  | Ala        | Leu        | Ser       |
| 374        | _     |            | 275 |             | _,         | _,     | _,         | 280   |            | _          | _    | _          | 285  | <b>~</b> 1 | _          | <b>.</b>  |
| 377<br>378 | Ser   | G1n<br>290 | Pro | He          | Phe        | Thr    | Thr<br>295 | Ala   | iie        | Asp        | Asp  | Arg        | Cys  | GIU        | ьeu        | ser       |
|            | Ser   |            | Arq | Glu         | Asp        | Thr    |            | Pro   | Ile        | Pro        | Ser  |            | Thr  | Ser        | Cys        | Thr       |
| 382        | 305   |            |     |             |            | 310    |            |       |            |            | 315  |            |      |            |            | 320       |
|            | Glu   | Thr        | Ser | Asp         |            | Leu    | Pro        | Thr   | Asn        |            | Asn  | Asp        | Asp  | Asp        |            | Cys       |
| 386        | Lvc   | Two        | Dro | Cvc         | 325        | Ual    | λla        | Pro   | Λen        | 330        | Tla  | Dro        | Leu  | Val        | 335<br>Ser | Ser       |
| 390        | пур   | цуъ        | PIO | 340         | 261        | Vai    | AIa        | FIO   | 345        | vob        | 116  | FIO        | Бец  | 350        | DCI        | DCI       |
|            | Thr   | Asn        | Leu | Ile         | Asn        | Glu    | Ile        | Asn   | Gly        | Val        | Ser  | Glu        | Lys  | Leu        | Ser        | Ala       |
| 394        |       |            | 355 |             |            |        |            | 360   | _          |            | ~-7  |            | 365  | _          | _          | <b></b> 1 |
| 397<br>398 | Thr   | Glu<br>370 | Ser | Ile         | Val        | Glu    | 11e<br>375 | Val   | Lys        | Gln        | GIu  | Val<br>380 | Leu  | Pro        | Leu        | Thr       |
|            | Leu   |            | Leu | Glu         | Ile        | Leu    |            | Asn   | Pro        | Pro        | Glu  |            | Met  | Lys        | Leu        | Glu       |
|            | 385   |            |     |             |            | 390    |            |       |            |            | 395  |            |      | •          |            | 400       |
|            | Cys   | Ile        | Pro | Ala         |            | Ile    | Thr        | Pro   | Ser        |            | Val  | Pro        | Ser  | Phe        |            | Pro       |
| 406        | Thr   | Dro        | Dro | Thr         | 405<br>Bro | Pro    | 71 =       | Sar   | Pro        | 410<br>Pro | Hic  | Thr        | Pro  | V=1        | 415        | Val       |
| 410        | 1111  | FIO        | FIO | 420         | FIO        | FIO    | AIG        | 261   | 425        | 110        | 1113 | 1111       | 110  | 430        | 110        | Vai       |
|            | Pro   | Ala        | Ala | Ala         | Thr        | Thr    | Val        | Ser   | Ser        | Pro        | Ser  | Ala        | Ala  | Ile        | Thr        | Val       |
| 414        |       |            | 435 | _           |            |        | _          | 440   | _          |            | _    | 1          | 445  | _          | _          | a l       |
| 417<br>418 | Gln   | Arg<br>450 | Val | Leu         | Glu        | Glu    | Asp<br>455 | Glu   | Ser        | He         | Arg  | Thr<br>460 | Cys  | Leu        | Ser        | GIu       |
|            | Asp   |            | Lvs | Glu         | Ile        | Gln    |            | Lvs   | Ile        | Glu        | Val  |            | Ala  | Asp        | Gly        | Gln       |
|            | 465   |            |     |             |            | 470    |            | 1     |            |            | 475  |            |      | •          | -          | 480       |
| 425        | Thr   | Glu        | Glu | Ile         |            | Asp    | Ser        | Gln   | Asn        |            | Asn  | Ser        | Arg  | Arg        |            | Pro       |
| 426        | 11-7  | D          | 77- | <b>G1</b> ~ | 485        | 77-    | 71.        | mb so | 770 J      | 490        | T    | mb ~       | Ten  | Tira       | 495        | Dro       |
| 429        | vaı   | Pro        | ATG | 500         | тте        | AIG    | тте        | Tur.  | va1<br>505 | PLO        | гу   | Int        | Trp  | ьуs<br>510 | пув        | PIO       |
|            | Lys   | Asp        | Arg |             | Arg        | Thr    | Thr        | Glu   |            | Met        | Leu  | Glu        | Ala  |            | Leu        | Glu       |
| 434        | _     | _          | 515 |             | _          |        |            | 520   |            |            |      |            | 525  |            |            |           |
|            | Leu   | -          | Ala | Glu         | Glu        | Glu    |            | Ser   | Ile        | Asp        | Lys  |            | Leu  | Glu        | Ser        | Glu       |
| 438        | Gln   | 530        | Lve | Met         | Ser        | Gln    | 535        | Phe   | Hic        | Pro        | Glu  | 540<br>Ara | Asp  | Pro        | Ser        | Asn       |
| -1-11      | GIII  | Tob        | пур | riec        | DET        | 0111   | CIY        | 1116  | 1113       | 110        | JIU  | A- 9       | 7.2P | 1+0        |            | - 12 P    |

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 04/26/2006 PATENT APPLICATION: US/10/574,398 TIME: 07:32:53

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## Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of/each sequence, which presents at least one n or Xaa.

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Seq#:7; N Pos. 377
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Seq#:41; N Pos. 1536
Seq#:43; N Pos. 34,115,321,356,359,378,385,394
Seg#:44; Xaa Pos. 12,39,107,119,120,129,132
Seq#:47; N Pos. 1871,1880,2479,2899
Seq#:48; Xaa Pos. 624,627,827,967
Seq#:65; N Pos. 477
Seq#:86; N Pos. 467,481,508,529,559
Seq#:88; N Pos. 305
Seq#:98; N Pos. 528,569,606
Seq#:106; N Pos. 639
Seq#:112; N Pos. 611,730,798,800
Seq#:142; N Pos. 464
Seq#:150; N Pos. 276,343,359,364,398,405,409
Seq#:152; N Pos. 179,203,503,505,506,508,568,627,629,680,694,702
Seq#:154; N Pos. 147,428,534,566,572
Seq#:160; N Pos. 522
Seq#:164; N Pos. 282
Seq#:188; N Pos. 149,155,156,157,160,180,201,203,396,411,454,475,495,514
Seq#:188; N Pos. 521,530,553,564,579,586,607,613
Seq#:222; N Pos. 383,392,416,477,483,523,564,593,602,619,627
Seq#:226; N Pos. 395
Seq#:238; N Pos. 797
Seq#:294; N Pos. 545,571,588,593,656
Seq#:314; N Pos. 569
Seq#:326; N Pos. 622
Seq#:336; N Pos. 370,426,458,471,483,484,510,519,535,538
Seq#:344; N Pos. 572,687
Seq#:361; N Pos. 47,271,290,333,359,466,467,582,655,669,895
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VERIFICATION SUMMARY

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Input Set : F:\5176.ST25.txt

Output Set: N:\CRF4\04262006\J574398.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:277 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:240 L:285 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:480 L:289 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:600 L:291 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:660 L:1067 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:360 L:1808 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:480 L:1810 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:540 L:3848 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:60 L:3852 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:180 L:3860 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:420 L:5704 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:1500 L:5829 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43 after pos.:0  $L:5831 \ M:341 \ W:$  (46) "n" or "Xaa" used, for SEQ ID#:43 after pos.:60 L:5839 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43 after pos.:300 L:5841 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43 after pos.:360 L:5884 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44 after pos.:0 L:5892 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44 after pos.:32 L:5908 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44 after pos.:96 L:5912 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44 after pos.:112 L:5916 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44 after pos.:128 L:6147 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:47 after pos.:1860 L:6167 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:47 after pos.:2460 L:6181 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:47 after pos.:2880 L:6378 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48 after pos.:608 L:6382 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48 after pos.:624 L:6430 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48 after pos.:816 L:6466 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48 after pos.:960 L:9048 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:65 after pos.:420 L:13011 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:86 after pos.:420 L:13013 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:86 after pos.:480 L:13015 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:86 after pos.:540 L:13127 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:88 after pos.:300 L:13975 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:98 after pos.:480 L:13977 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:98 after pos.:540 L:13979 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:98 after pos.:600 L:16229 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:106 after pos.:600 L:16510 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:112 after pos.:600 L:16514 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:112 after pos.:720 L:16516 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:112 after pos.:780 L:19326 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:142 after pos.:420 L:19922 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:150 after pos.:240 L:19924 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:150 after pos.:300 L:19926 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:150 after pos.:360 L:20084 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:152 after pos.:120 L:20086 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:152 after pos.:180 L:20096 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:152 after pos.:480

VERIFICATION SUMMARY DATE: 04/26/2006
PATENT APPLICATION: US/10/574,398 TIME: 07:32:53

Input Set : F:\5176.ST25.txt

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L:20098 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:152 after pos.:540
L:20100 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:152 after pos.:600
L:20102 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:152 after pos.:660
L:20404 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:154 after pos.:120
L:20414 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:154 after pos.:420
L:20416 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:154 after pos.:480
L:20418 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:154 after pos.:540
L:20826 \ M:341 \ W: (46) "n" or "Xaa" used, for SEQ ID#:160 after pos.:480
L:21296 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:164 after pos.:240
L:24245 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:188 after pos.:120
L:24247 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:188 after pos.:180
L:24253 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:188 after pos.:360
L:24255 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:188 after pos.:420
L:24257 \ M:341 \ W: (46) "n" or "Xaa" used, for SEQ ID#:188 after pos.:480
L:24259 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:188 after pos.:540
L:24261 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:188 after pos.:600
L:27719 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:222 after pos.:360
L:27721 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:222 after pos.:420
L:27723 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:222 after pos.:480
L:27725 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:222 after pos.:540
L:27727 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:222 after pos.:600
L:28439 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:226 after pos.:360
L:30299 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:238 after pos.:780
L:39805 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:294 after pos.:540
L:39807 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:294 after pos.:600
L:42885 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:314 after pos.:540
L:44337 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:326 after pos.:600
L:45307 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:336 after pos.:360
L:45309 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:336 after pos.:420
L:45311 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:336 after pos.:480
L:45942 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:344 after pos.:540
L:45946 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:344 after pos.:660
L:47542 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:361 after pos.:0
L:47550 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:361 after pos.:240
L:47552 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:361 after pos.:300
L:47556 M:341 W: (46)
                     "n" or "Xaa" used, for SEQ ID#:361 after pos.:420
L:47560 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:361 after pos.:540
L:47562 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:361 after pos.:600
L:47564 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:361 after pos.:660
L:47570 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:361 after pos.:840
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